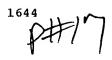


AUG 02 2000

TECH CENTER 1600/2900



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/121,017B

Input Set : A:\revised sequence listing.txt Output Set: N:\CRF3\07142000\I121017B.raw

DATE: 07/14/2000

TIME: 10:22:52

ENTERE

```
5 <110> APPLICANT: Imamura, Toru
         Asada, Masahiro
 9
         Oka, Syuichi
11
         Suzuki, Masashi
         Yoneda, Atsuko
15
         Ota, Keiko
         Oda, Yuko
         Miyakawa, Kazuko
19
         Orikasa, Noriko
21
         Asada, Chie
23
         Kojima, Tetsuhito
29 <120> TITLE OF INVENTION: HEPARIN-BINDING PROTEINS MODIFIED WITH SUGAR CHAINS,
31
         METHOD OF PRODUCING THE SAME AND PHARMACEUTICAL
33
         COMPOSITIONS CONTAINING THE SAME
37 <130> FILE REFERENCE: 382.1019
41 <140> CURRENT APPLICATION NUMBER: 09/121,017B
43 <141> CURRENT FILING DATE: 1998-07-22
47 <150> PRIOR APPLICATION NUMBER: 307721/1997
49 <151> PRIOR FILING DATE: 1997-11-10
53 <160> NUMBER OF SEO ID NOS: 31
57 <170> SOFTWARE: PatentIn Ver. 2.0
61 <210> SEQ ID NO: 1
63 <211> LENGTH: 221
65 <212> TYPE: PRT
67 <213> ORGANISM: Artificial Sequence
71 <220> FEATURE:
73 <223> OTHER INFORMATION: Description of Artificial Sequence: fusion of
75
         sequence for a part of human rydocan and a part of human fibroblast
77
         growth factor 1
81 <400> SEQUENCE: 1
83 Met Ala Pro Ala Arg Leu Phe Ala Leu Leu Phe Phe Val Gly Gly
89 Val Ala Glu Ser Ile Arg Glu Thr Glu Val Ile Asp Pro Gln Asp Leu
                20
                                    25
95 Leu Glu Gly Arg Tyr Phe Ser Gly Ala Leu Pro Asp Asp Glu Asp Val
                                40
101 Val Gly Pro Gly Gln Glu Ser Asp Asp Phe Glu Leu Ser Gly Ser Gly
                             55
107 Asp Leu Asp Asp Leu Glu Asp Ser Met Ile Gly Pro Glu Val Val His
113 Pro Leu Val Pro Leu Asp Ala Asn Tyr Lys Lys Pro Lys Leu Leu Tyr
119 Cys Ser Asn Gly Gly His Phe Leu Arg Ile Leu Pro Asp Gly Thr Val
121 100 105 110
125 Asp Gly Thr Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln Leu Ser
                                120
           115
                                                     125
131 Ala Glu Ser Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr Gly Gln
```

~~ 02 2000

Concentrate all'isso

RECEIVED

Input Set : A:\revised sequence listing.txt
Output Set: N:\CRF3\07142000\I121017B.raw

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/121,017B

DATE: 07/14/2000 TIME: 10:22:52

133		130					135					140					
137	Tyr	Leu	Ala	Met	Asp	Thr	Asp	Gly	Leu	Leu		Gly	Ser	Gln	Thr	Pro	
139	145					150					155					160	
143	Asn	Glu	Glu	Cys	Leu	Phe	Leu	Glu	Arg	Leu	Glu	Glu	Asn	His	Tyr	Asn	
145					165					170					175		
149	Thr	Tyr	Ile	Ser	Lys	Lys	His	Ala	Glu	Lys	Asn	Trp	Phe	Val	Gly	Leu	
151	•			180					185					190			
155	Lys	Lys	Asn	Gly	Ser	Cys	Lys	Arg	Gly	Pro	Arg	Thr	His	Tyr	Gly	Gln	
157			195					200					205				
	Lys		Ile	Leu	Phe	Leu		Leu	Pro	Val	Ser		Asp				
163		210					215					220					
	<210> SEQ ID NO: 2																
	<211> LENGTH: 663																
	<212> TYPE: DNA																
	<213> ORGANISM: Artificial Sequence																
	<pre><220> FEATURE:</pre>																
	<22.															fusio	
183							rt o	t hur	nan 1	rydo	can a	and a	a par	rt o	t hur	nan fil	problast
185		_	cowtl		ctor	1											
			EATUE		050												
			AME/I														
			CAT			(6 t	33)										
			EQUE				++-	~~~	a+ a	-+-		++~	++-	~+ ~		~~~	4.0
	_	•	ccc Pro	-	-	-			-	-	-			-			48
201	1	ніа	PIO	HIG	5	neu	FIIC	Ala	ьец	10	пеа	File	riie	vai	15	Gry	
	_	acc	gag	tca	_	cas	αаσ	act	дад		atc	gac	CCC	cad		ctc	96
			Glu														50
211				20					25					30		200	
	cta	gaa	ggc		tac	ttc	tcc	qqa	qcc	cta	cca	gac	gat	gag	gat	qta	144
			Gly														
219			35	_	-			40				•	45		-		
223	gtg	ggg	ccc	ggg	cag	gaa	tct	gat	gac	ttt	gag	ctg	tct	ggc	tct	gga	192
225	Val	Gly	Pro	Gly	Gln	Glu	Ser	Asp	Asp	Phe	Glu	Leu	Ser	Gly	Ser	Gly	
227		50					55					60					
231	gat	ctg	gat	gac	ttg	gaa	gac	tcc	atg	atc	ggc	cct	gaa	gtt	gtc.	cat	240
		Leu	Asp	Asp	Leu	Glu	Asp	Ser	Met	Ile	Gly	Pro	Glu	Val	Val	His	
	65					70					75			1		80	
			gtg														288
	Pro	Leu	Val	Pro		Asp	Ala	Asn	Tyr	7	Lys	Pro	Lys	Leu		Tyr	
243					. 85					90					95		
			aac														336
	Cys	ser	Asn		GTA	HIS	rne	ren		rre	ьeu	Pro	ASP		Thr	vaı	
251				100					105					110			304
			aca														384
259	изр	стХ	Thr 115	Arg	Asp	AIG	ser	120	GIII	utz	tre	GTU		OTII	ьeu	ser	
	~~~	<i>a</i>		a+~	~~~	~ - ~	a+~		a+=		a a t	300	125	3.0±	~~~	020	432
			agc Ser														13 3 4
203	urd	GIU	261	va⊥	GIY	GIU	4 a I	TYL	116	пур	261	1111	GIU	TILL	оту	GIII	

## RAW SEQUENCE LISTING DATE: 07/14/2000 PATENT APPLICATION: US/09/121,017B TIME: 10:22:52

AUG 02 2000

RECEIVED

Input Set : A:\revised sequence listing.txt
Output Set: N:\CRF3\07142000\I121017B.raw

TECH CENTER IF

```
267
                                   135
271 tac ttg gcc atg gac acc gac ggg ctt tta tac ggc tca cag aca cca
273 Tyr Leu Ala Met Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln Thr Pro
275 145 150 155 160
279 aat gag gaa tgt ttg ttc ctg gaa agg ctg gag gag aac cat tac aac 281 Asn Glu Glu Cys Leu Phe Leu Glu Arg Leu Glu Glu Asn His Tyr Asn 283 165 170 175
                                                                                        528
                                                                                        576
287 acc tat ata tcc aag aag cat gca gag aag aat tgg ttt gtt ggc ctc
289 Thr Tyr Ile Ser Lys Lys His Ala Glu Lys Asn Trp Phe Val Gly Leu
291
                  180
                                            185
295 aag aag aat ggg agc tgc aaa cgc ggt cct cgg act cac tat ggc cag
297 Lys Lys Asn Gly Ser Cys Lys Arg Gly Pro Arg Thr His Tyr Gly Gln
299 195
                                     200
303 aaa gca atc ttg ttt ctc ccc ctg cca gtc tct tct gat
305 Lys Ala Ile Leu Phe Leu Pro Leu Pro Val Ser Ser Asp
307 210 215 220
                                                                                        663
313 <210> SEQ ID NO: 3
315 <211> LENGTH: 175
317 <212> TYPE: PRT
319 <213> ORGANISM: Artificial Sequence
323 <220> FEATURE:
325 <223> OTHER INFORMATION: Description of Artificial Sequence: fusion of
327
            sequence for a part of mouse fibroblast growth factor 6 and
329
            a part of human fibroblast growth factor 1
333 <400> SEQUENCE: 3
335 Met Ser Arg Gly Ala Gly Arg Val Gln Gly Thr Leu Gln Ala Leu Val 337 1 5 10 15
341 Phe Leu Gly Val Leu Val Gly Met Val Val Pro Ser Pro Ala Gly Ala 343 20 25 30
347 Arg Ala Asn Gly Thr Leu Leu Asp Ala Asn Tyr Lys Lys Pro Lys Leu 349 35 40 45
353 Leu Tyr Cys Ser Asn Gly Gly His Phe Leu Arg Ile Leu Pro Asp Gly 355 \phantom{000}55\phantom{000} 60
359 Thr Val Asp Gly Thr Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln
361 65 70 75 80
365 Leu Ser Ala Glu Ser Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr
367 85 90 95
371 Gly Gln Tyr Leu Ala Met Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln 373 100 \, 105 110 \,
377 Thr Pro Asn Glu Glu Cys Leu Phe Leu Glu Arg Leu Glu Glu Asn His
379 115 120 125
383 Tyr Asn Thr Tyr Ile Ser Lys Lys His Ala Glu Lys Asn Trp Phe Val
385 130 135 140
389 Gly Leu Lys Lys Asn Gly Ser Cys Lys Arg Gly Pro Arg Thr His Tyr 391 145 150 150 155
395 Gly Gln Lys Ala Ile Leu Phe Leu Pro Leu Pro Val Ser Ser Asp
401 <210> SEQ ID NO: 4
403 <211> LENGTH: 525
```

RAW SEQUENCE LISTING DATE: 07/14/2000 PATENT APPLICATION: US/09/121,017B TIME: 10:22:52

Input Set: A:\revised sequence listing.txt
Output Set: N:\CRF3\07142000\I121017B.raw

```
405 <212> TYPE: DNA
407 <213> ORGANISM: Artificial Sequence
411 <220> FEATURE:
413 <223> OTHER INFORMATION: Description of Artificial Sequence: fusion of
            sequence for a part of mouse fibroblast growth factor 6 and
415
            a part of human fibroblast growth factor 1
417
421 <220> FEATURE:
423 <221> NAME/KEY: CDS
425 <222> LOCATION: (1)..(525)
429 <400> SEQUENCE: 4
431 atg tcc cgg gga gca gga cgt gtt cag ggc acg ctg cag gct ctc gtc
433 Met Ser Arg Gly Ala Gly Arg Val Gln Gly Thr Leu Gln Ala Leu Val
435 1
                          5
                                                 10
439 ttc tta ggc gtc cta gtg ggc atg gtg gtg ccc tca cct gcc ggc gcc 441 Phe Leu Gly Val Leu Val Gly Met Val Val Pro Ser Pro Ala Gly Ala 443 20 25 30
447 cgc gcc aac ggc acg cta ctg gac gct aat tac aag aag ccc aaa ctc
449 Arg Ala Asn Gly Thr Leu Leu Asp Ala Asn Tyr Lys Lys Pro Lys Leu
451
               35
                                    40
455 ctc tac tgt agc aac ggg ggc cac ttc ctg agg atc ctt ccg gat ggc
457 Leu Tyr Cys Ser Asn Gly Gly His Phe Leu Arg Ile Leu Pro Asp Gly
    50
                                 55
463 aca gtg gat ggg aca agg gac agg agc cag cac att cag ctg cag 465 Thr Val Asp Gly Thr Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln 467 65 70 75 80
471 ctc agt gcg gaa agc gtg ggg gag gtg tat ata aag agt acc gag act 473 Leu Ser Ala Glu Ser Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr 475 85 90 95
                                                                                        288
479 ggc cag tac ttg gcc atg gac acc gac ggg ctt tta tac ggc tca cag 481 Gly Gln Tyr Leu Ala Met Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln
                                                                                        336
483
                   100
                                            105
                                                                     110
487 aca cca aat gag gaa tgt ttg ttc ctg gaa agg ctg gag gag aac cat
489 Thr Pro Asn Glu Glu Cys Leu Phe Leu Glu Arg Leu Glu Glu Asn His
491 115 120 125
495 tac aac acc tat ata tcc aag aag cat gca gag aag aat tgg ttt gtt
497 Tyr Asn Thr Tyr Ile Ser Lys Lys His Ala Glu Lys Asn Trp Phe Val
499 130 135 140
503 ggc ctc aag aag aat ggg agc tgc aaa cgc ggt cct cgg act cac tat 505 Gly Leu Lys Lys Asn Gly Ser Cys Lys Arg Gly Pro Arg Thr His Tyr
                                                                                        480
507 145
                             150
                                                      155
511 ggc cag aaa gca atc ttg ttt ctc ccc ctg cca gtc tct tct gat
513 Gly Gln Lys Ala Ile Leu Phe Leu Pro Leu Pro Val Ser Ser Asp
                                                                                        525
515
                        165
                                                 170
523 <210> SEQ ID NO: 5
525 <211> LENGTH: 181
527 <212> TYPE: PRT
529 <213> ORGANISM: Artificial Sequence
533 <220> FEATURE:
535 <223> OTHER INFORMATION: Description of Artificial Sequence: fusion of
```

RAW SEQUENCE LISTING DATE: 07/14/2000 PATENT APPLICATION: US/09/121,017B TIME: 10:22:52

Input Set : A:\revised sequence listing.txt
Output Set: N:\CRF3\07142000\I121017B.raw

```
537
           sequence for a part of mouse fibroblast growth factor 6,
539
           a part of human fibroblast growth factor 1 and an artificial
541
           sequence
545 <400> SEQUENCE: 5
547 Met Ser Arg Gly Ala Gly Arg Val Gln Gly Thr Leu Gln Ala Leu Val
549 1 5 10 15
559 Arg Ala Gln Gly Thr Leu Leu Asp Ala Asn Tyr Lys Lys Pro Lys Leu
561 35 40 45
565 Leu Tyr Cys Ser Asn Gly Gly His Phe Leu Arg Ile Leu Pro Asp Gly 567 50 60
571 Thr Val Asp Gly Thr Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln 573 65 70 75 80
577 Leu Ser Ala Glu Ser Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr 579 85 90 95
583 Gly Gln Tyr Leu Ala Met Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln 585 \phantom{\bigg|}100\phantom{\bigg|}105\phantom{\bigg|}
589 Thr Pro Asn Glu Glu Cys Leu Phe Leu Glu Arg Leu Glu Glu Ala Ala 591 $125$
595 Thr Pro Ala Pro Asn His Tyr Asn Thr Tyr Ile Ser Lys Lys His Ala
597 130 135 140
601 Glu Lys Asn Trp Phe Val Gly Leu Lys Lys Asn Gly Ser Cys Lys Arg
603 145 150 155 160
607 Gly Pro Arg Thr His Tyr Gly Gln Lys Ala Ile Leu Phe Leu Pro Leu
                    165
613 Pro Val Ser Ser Asp
615
                 180
621 <210> SEQ ID NO: 6
623 <211> LENGTH: 543
625 <212> TYPE: DNA
627 <213> ORGANISM: Artificial Sequence
631 <220> FEATURE:
633 <223> OTHER INFORMATION: Description of Artificial Sequence: fusion of
           sequence for a part of mouse fibroblast growth factor 6,
           a part of human fibroblast growth factor 1 and an artificial
639
          sequence
643 <220> FEATURE:
645 <221> NAME/KEY: CDS
647 <222> LOCATION: (1)..(543)
651 <400> SEQUENCE: 6
653 atg tee egg gga gea gga egt gtt eag gge aeg etg eag get ete gte
                                                                                48
655 Met Ser Arg Gly Ala Gly Arg Val Gln Gly Thr Leu Gln Ala Leu Val
657 1
                       5
                                              10
661 ttc tta ggc gtc cta gtg ggc atg gtg gtg ccc tca cct gcc ggc gcc 663 Phe Leu Gly Val Leu Val Gly Met Val Val Pro Ser Pro Ala Gly Ala 665 20 \hspace{1.5cm} 20 \hspace{1.5cm} 30
669 cgc gcc caa ggc acg cta ctg gac gct aat tac aag aag ccc aaa ctc
671 Arg Ala Gln Gly Thr Leu Leu Asp Ala Asn Tyr Lys Lys Pro Lys Leu
```

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/121,017B

DATE: 07/14/2000 TIME: 10:22:53

Input Set : A:\revised sequence listing.txt
Output Set: N:\CRF3\07142000\I121017B.raw

L:1763 M:258 W: Mandatory Feature missing, <220> FEATURE: